

## Lab Resource: Stem Cell Line

# Generation of an induced pluripotent stem cell line (CSC-44) from a Parkinson's disease patient carrying a compound heterozygous mutation (c.823C>T and EX6 del) in the *PARK2* gene



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## ABSTRACT

Mutations in the *PARK2* gene, which encodes PARKIN, are the most frequent cause of autosomal recessive Parkinson's disease (PD). We report the generation of an induced pluripotent stem cell (iPSC) line from a 78-year-old patient carrying a compound heterozygous mutation (c.823C>T and EX6del) in the *PARK2* gene. Skin fibroblasts were reprogrammed using the non-integrating Sendai virus technology to deliver OCT3/4, SOX2, c-MYC and KLF4 factors. The generated cell line CSC-44 exhibits expression of common pluripotency markers, *in vitro* differentiation into the three germ layers and normal karyotype. This iPSC line can be used to explore the association between *PARK2* mutations and PD.

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## Resource table.

Unique stem cell line identifier	ULUNDi006-A
Alternative name(s) of stem cell line	CSC-44i
Institution	Stem Cell Laboratory for CNS Disease Modeling, Department of Experimental Medical Science, Lund University
Contact information of distributor	Laurent Roybon, <a href="mailto:Laurent.Roybon@med.lu.se">Laurent.Roybon@med.lu.se</a>
Type of cell line	iPSC
Origin	Human
Additional origin info	Age of patient at onset: 33 Sex of patient: female Ethnicity: N/A
Cell source	Skin fibroblasts
Clonality	Clonal
Method of reprogramming	Sendai virus mediated delivery of OCT3/4, SOX2, c-MYC and KLF4

## (continued)

Genetic modification	No modification
Type of modification	No modification
Associated disease	Parkinson's disease
Gene/locus	<i>PARK2</i> (MIM #602544) on chromosome 6q26 Genotype: compound heterozygous mutation: c.823C>T in exon 7 and deletion of exon 6
Method of modification	No modification
Name of transgene or resistance	No transgene or resistance
Inducible/constitutive system	Not inducible
Date archived/stock date	N/A
Cell line repository/bank	N/A
Ethical approval	Parkinson Institute Biobank (part of the Telethon Genetic Biobank Network <a href="http://biobanknetwork.telethon.it/">http://biobanknetwork.telethon.it/</a> ): approved by Ethics Committee "Milano Area C" ( <a href="http://comitatoeticoareac.ospedaleniguarda.it/">http://comitatoeticoareac.ospedaleniguarda.it/</a> ) on the 26/06/2015, Numero Registro dei pareri: 370-062015. Reprogramming: 202100-3211 (delivered by Swedish work environment Arbetsmiljöverket).

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## Resource utility

Mutations in the *PARK2* gene are the second most common known cause of Parkinson's disease (PD). CSC-44 iPSC line was generated from a PD patient with a compound heterozygous mutation in *PARK2*. This iPSC line can be used as a model to explore the link between mutant *PARK2* and PD pathology.

## Resource details

Mutations in *PARK2* result in autosomal-recessive familial PD and are the second most common known cause of this neurodegenerative disease. *PARK2* encodes for PARKIN, a E3 ubiquitin ligase that plays a role in targeting proteins for degradation and maintaining mitochondrial function (Nuytemans et al., 2010). Among the identified *PARK2* mutations, both deletions and insertions of one or more exons and missense mutations have been described. Here, we report the generation of an induced pluripotent stem cell line (CSC-44) from a patient with PD caused by a compound heterozygous *PARK2* mutation: point mutation c.823C>T in exon 7 and deletion of exon 6. The c.823C>T mutation in *PARK2* gene predicts an arginine to tryptophan substitution at amino acid residue 275 (p.R275W), which is located within the RING1 domain of PARKIN. This mutation disrupts the charge distribution and leads to local rearrangements in the RING1-IBR interface, which hampers ubiquitin ligase activity and confers a toxic gain of function to PARKIN, leading to its aggregation (Fiesel et al., 2015; Oczkowska et al., 2013). Even though this point mutation has been well characterized and associated with other mutations in compound heterozygotes, its association with the deletion of exon 6 has not been described. Given the importance of compound heterozygous mutations on *PARK2* gene, CSC-44 iPSC line can be used to better understand the impact of both alterations on cellular function.

To generate this line, skin fibroblasts collected by punch skin biopsy from a 78-year-old PD patient were reprogrammed using a non-integrating Sendai virus technology. Briefly, fibroblasts were seeded (75,000 cells/well) on a 12-well plate, two days before transduction. The CytoTune™-iPS 2.0 Sendai Reprogramming Kit was then used to deliver the four reprogramming factors (OCT3/4, SOX2, c-MYC and KLF4). At day 7 post-transduction, the cells were re-seeded onto mouse embryonic fibroblasts (MEF)-feeder layer and expanded until colonies presented an embryonic stem cell (ES)-like morphology. At day 28, 12 colonies were picked and expanded as individual clones for 7 days. Three clones (CSC-44G, CSC-44I, CSC-44J) were further selected for expansion and karyotype analysis. All clones were characterized using the methods we previously described (Holmqvist et al., 2016). Here, we present the characterization of clone CSC-44I.

The generated clone, CSC-44I, expressed the common nuclear and cell surface pluripotency markers, OCT4/NANOG and TRA1-81/SSEA4 (Fig. 1A) and showed alkaline phosphatase (ALP) activity (Fig. 1B). Flow cytometry analysis demonstrated that >98% of the iPSCs were positive for SSEA4 (Fig. 1C). Additional immunocytochemistry analysis revealed elimination of the Sendai virus at passage 7 (Fig. 1D). Fig. 1E presents a normal female karyogram identified in CSC-44I clone. The identity of the generated iPSC line was confirmed by DNA fingerprint, showing genetic correspondence to parental fibroblasts. DNA sequencing analysis of CSC-44I iPSCs confirmed the presence of mutations in the *PARK2* gene (Fig. 1F). Embryoid bodies (EBs) generated from CSC-44I iPSCs present three-germ layer differentiation capacity as shown by the *in vitro* expression of alpha-fetoprotein (AFP), an endodermal marker, smooth muscle actin (SMA), a mesodermal marker, and beta-III-tubulin (BIIIITub), an ectodermal marker (Fig. 1G). Mycoplasma infection was prevented by routine addition of plasmocin in cell culture media at early passages.

## Materials and methods

### Fibroblast culture

Dermal fibroblasts were collected by punch skin biopsy from a patient diagnosed with PD, after obtaining informed consent. The fibroblasts were maintained in fibroblast growth medium, composed of DMEM media (ThermoFisher Scientific) with 10% fetal bovine serum and 1% Penicillin-Streptomycin and passaged with 0.05% trypsin.

### iPSC generation and expansion

For reprogramming, 75,000 cells were seeded on a 12-well plate and maintained in fibroblast growth medium. Two days after (day 0), the cells were transduced using the three vector preparations (MOI = 5, 5, 3) included in the CytoTune™-iPS 2.0 Sendai Reprogramming Kit (ThermoFisher Scientific). On the following day and on every other day, the medium was replaced with fresh fibroblast growth medium. At day 7, the cells were re-seeded onto irradiated mouse embryonic fibroblasts (MEFs) feeder cells with fibroblast growth medium. On the day after and until colony picking, the cells were cultured in WiCell medium composed of advanced DMEM/F12 (ThermoFisher Scientific), 20% Knock-Out Serum Replacement (v/v, ThermoFisher Scientific), 2 mM L-glutamine (ThermoFisher Scientific), 1% non-essential amino acids (NEAA, v/v, Millipore) and 0.1 mM  $\beta$ -mercaptoethanol (Sigma-Aldrich), supplemented with 20 ng/ml FGF2 (ThermoFisher Scientific). On day 28, individual colonies were picked and re-seeded on a 24-well plate containing fresh MEFs. One week after, three clones were selected based on the morphology of the colonies, and further expanded on 6-well plates. The cells were passaged once a week and seeded on the appropriate cell culture surface for characterization assays at the indicated passage numbers (Table 1).

### Immunocytochemistry

The iPSCs cultures were fixed with 4% paraformaldehyde for 15 min at room temperature (RT), permeabilized and blocked for 1 h at RT with PBS containing 10% donkey serum and 0.1% TritonX-100 (Sigma) and incubated overnight at +4 °C with the primary antibodies (Table 2) diluted in the blocking buffer. The secondary antibodies were thereafter added for 1 h at RT in the dark, followed by nuclei counterstain with DAPI (1:10,000 (Life Technologies)) and image acquisition on inverted epifluorescence microscope LRI - Olympus IX-73. Scale bars are 200  $\mu$ m.

### Alkaline phosphatase activity

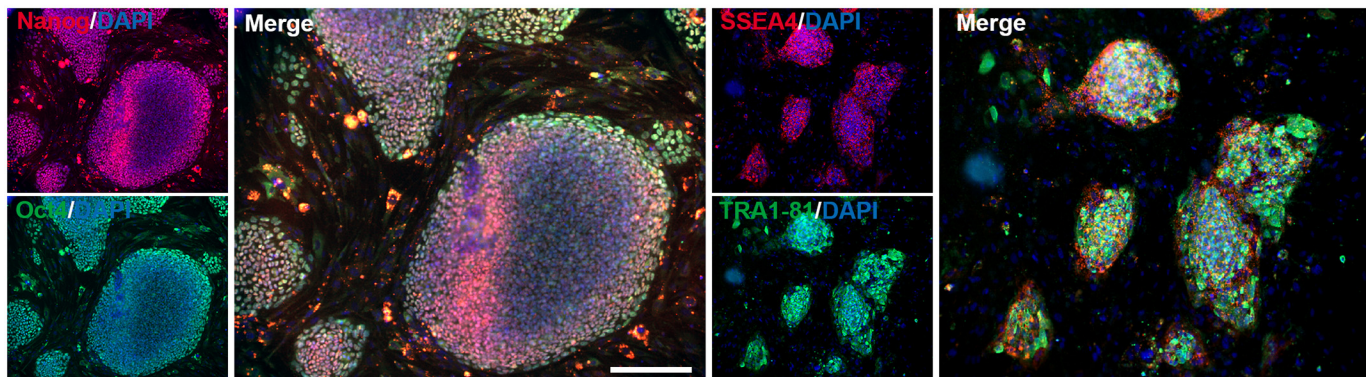
Alkaline phosphatase staining was performed using Alkaline Phosphatase Staining Kit (Stemgent, MA).

### *In vitro* differentiation by embryoid body (EB) formation

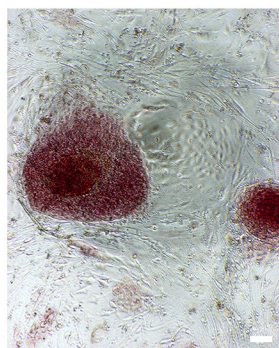
Human iPSCs were grown for 2 weeks as embryoid bodies (EBs) in low-attachment 24-well plates (Corning) in WiCell medium supplemented with 20 ng/ml FGF2. The EBs were then seeded on a 0.1% gelatin-coated 96-well plate (Greiner Bio-One) in DMEM media containing 10% fetal bovine serum and 1% Penicillin-Streptomycin for subsequent spontaneous differentiation, with media changes every 2–3 days. After 2 weeks, the cells were fixed and stained for three germ-layer markers as described in Table 2.



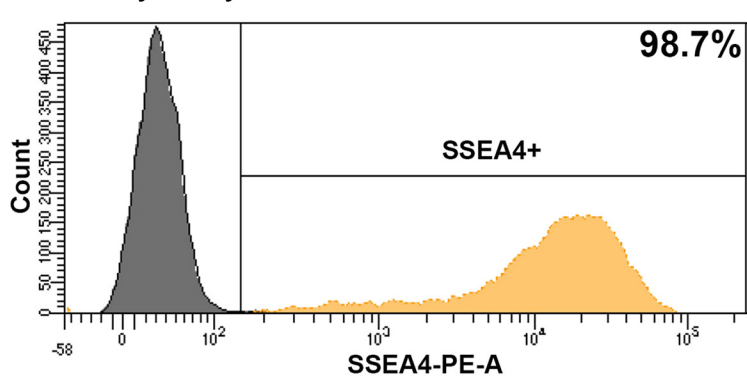
### A. Pluripotency markers



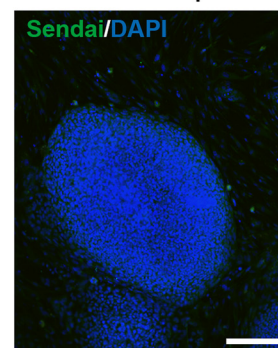
### B. Alkaline phosphatase



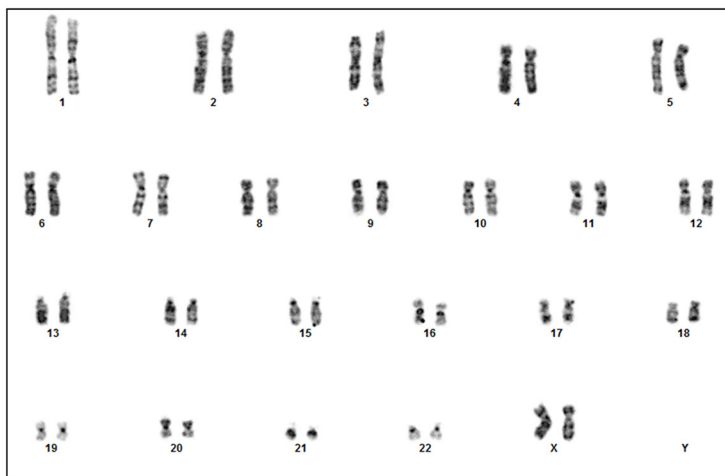
### C. Flow cytometry



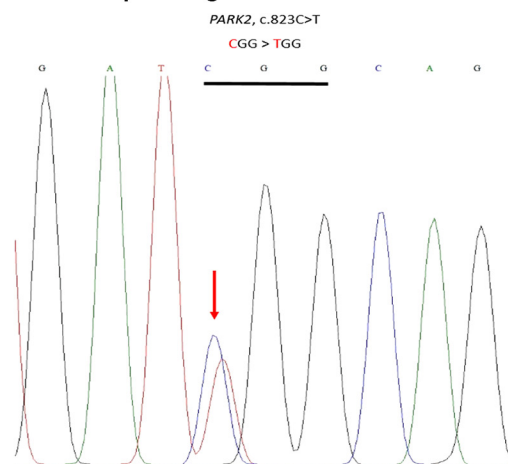
### D. Sendai virus expression



### E. Karyogram



### F. DNA Sequencing



### G. In vitro differentiation

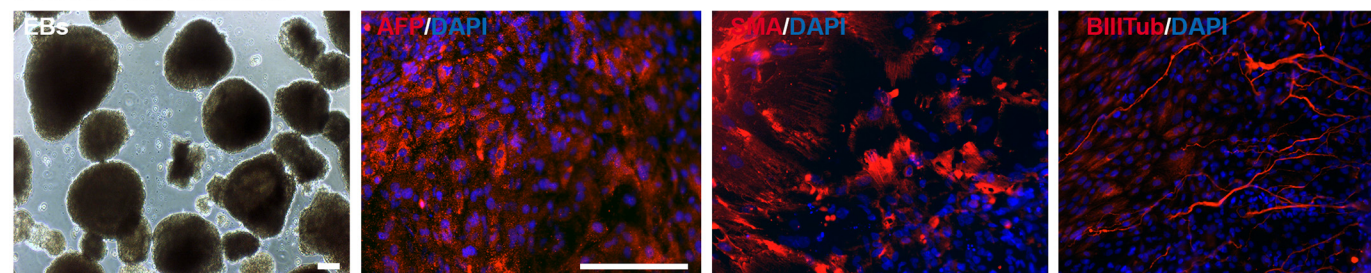


Fig. 1. Characterization of the iPSC line CSC-44I.

**Table 1**  
Characterization and validation.

Classification	Test	Result	Data
Morphology	Photography	Visual record of the line: normal	Not shown but available with author
Phenotype	Immunocytochemistry	Positive staining for pluripotency markers: OCT4, NANOG, TRA1-81 and SSEA4	Fig. 1 panel A
	Alkaline phosphatase activity	Visible activity	Fig. 1 panel B
	Flow cytometry	94.8% SSEA4	Fig. 1 panel C
Karyotype	G-banding	46,XX, (300–400 bands resolution in average)	Fig. 1 panel E
Identity	STR analysis	10 sites analyzed, all matched with parent fibroblast cell line	Available with author
Mutation analysis (IF APPLICABLE)	Sequencing	Compound heterozygous mutation in the <i>PARK2</i> gene (c.823C>T and EX6del)	Fig. 1 panel F
Microbiology and virology	Mycoplasma	Mycoplasma testing by RT-PCR. Negative.	Not shown but available with author
Differentiation potential	Embryoid body formation	Staining of smooth muscle actin, beta-III-tubulin and alpha-fetoprotein after spontaneous differentiation of embryoid bodies	Fig. 1 panel G
Donor screening (OPTIONAL)	HIV 1 + 2 Hepatitis B, Hepatitis C	N/A	N/A
Genotype additional info (OPTIONAL)	Blood group genotyping	N/A	N/A
	HLA tissue typing	N/A	N/A

### Karyotype analysis

The G-banding analysis was performed at 300–400 band resolution in average after 9 passages, at the Department of Clinical Genetics and Pathology in Lund.

### Mutation sequencing

Genomic DNA from fibroblasts and iPSCs was extracted using conventional lysis buffer composed of 100 mM Tris (pH 8.0), 200 mM NaCl, 5 mM EDTA and 0.2% SDS in distilled autoclaved water supplemented with 1.5 mg/ml Proteinase K. The mutation in the *PARK2* gene was confirmed by direct DNA sequencing (Macrogen Europe, Amsterdam, The Netherlands). Primers used for amplification and directed sequencing of *PARK2* around the mutation sites are listed in Table 2.

### DNA fingerprinting

Genomic DNA from fibroblasts and iPSCs was isolated as described above and fingerprinting analyses was outsourced to the IdentiCell STR profiling service (Department of Molecular Medicine, Aarhus University Hospital, Skejby, Denmark).

### Mycoplasma detection

Absence of mycoplasma contamination was confirmed by the real-time PCR method at GATC Biotech AG (European Genome and Diagnostics Centre, Konstanz, Germany).

### Acknowledgements

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**Table 2**  
Reagents details.

Antibodies used for immunocytochemistry			
	Antibody	Dilution	Company Cat # and RRID
Pluripotency markers	Mouse anti-Oct4	1:200	Millipore Cat# MAB4401, RRID:AB_2167852
	PE-conjugated mouse anti-human NANOG	1:200	BD Biosciences Cat# 560483, RRID:AB_1645522
	Mouse anti- TRA-1-81	1:200	Thermo Fisher Scientific Cat# 41-1100, RRID:AB_2533495
	PE-conjugated mouse anti-SSEA4	1:200	Thermo Fisher Scientific Cat# A14766, RRID:AB_2534281
Sendai	Chicken anti-Sendai virus	1:1000	Abcam Cat# ab33988, RRID:AB_777877
Differentiation markers	Mouse anti-AFP	1:200	Sigma-Aldrich Cat# A8452, RRID:AB_258392
	Mouse anti-SMA	1:200	Sigma-Aldrich Cat# A2547, RRID:AB_476701
	Mouse anti-beta-III tubulin	1:200	Sigma-Aldrich Cat# T8660, RRID:AB_477590
Secondary antibodies	Donkey anti-mouse Alexa Fluor® 488	1:400	Molecular Probes Cat# A-21202, RRID:AB_141607
	Donkey anti-chicken Alexa Fluor® 488	1:400	Jackson ImmunoResearch Labs Cat# 703-545-155, RRID:AB_2340375
	Donkey anti-mouse Alexa Fluor® 555	1:400	Thermo Fisher Scientific Cat# A-31570, RRID:AB_2536180
Primers			
	Target	Forward/reverse primer (5'-3')	
Mutation sequencing	<i>PARK2</i>	AGGATTACAGAAATTGGTCT/TCTGTCTTCATTAGCATTAGA	

## Appendix A. Supplementary data

Supplementary data to this article can be found online at <https://doi.org/10.1016/j.scr.2018.01.006>.

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